

**LIST OF PATENTS AND OTHER ITEMS FOR APPLICANT'S  
INFORMATION DISCLOSURE STATEMENT**

(Use several sheets if necessary)

00801.0197.NPUS00

09/864,637

**APPLICANT:**

Chia-Lin WEI, et al.

**FILING DATE:**

May 23, 2001

**GROUP:**

To be Assigned 1637

**U.S. PATENT DOCUMENTS**

EXAMINER INITIAL		DOCUMENT NUMBER	DATE	NAME	CLASS	SUB CLASS	FILING DATE
TS	✓ AA	4,366,241	12/28/82	Tom et al.	435	7	08/07/80
	✓ AB	B1 4,366,241	10/18/88	Tom et al. (Reexamination)	435	7	03/04/87
	✓ AC	4,582,789	04/15/86	Sheldon III et al.	435	6	12/18/84
	✓ AD	5,143,854	09/01/92	Pirrung et al.	436	518	03/07/90
	✓ AE	5,702,898	12/30/97	Bonaldo et al.	435	6	06/06/95
TS	✓ AF	5,846,721	12/8/98	Soares et al.	435	6	09/19/96

**FOREIGN PATENT DOCUMENTS**

EXAMINER INITIAL		DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUB CLASS	TRANSLATION YES NO
				None			

**OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages, etc.)**

TS	✓ AG	Bonaldo, Maria de Fatima, et al.; <i>Normalization and Subtraction: Two Approaches to Facilitate Gene Discovery</i> ; <b>Genome Research</b> ; Volume 6, No. 9; pages 791-806; (September, 1996).
I	✓ AH	Carninci, Piero, <i>Normalization and Subtraction of Cap-Trapper-Selected cDNAs to Prepare Full-Length cDNA Libraries for Rapid Discovery of New Genes</i> ; <b>Research</b> ; Vol 10, No. 10; pages 1617-1630; (October, 2000).
TS	✓ AI	Grimwade, David, et al.; <i>RT-PCR in Diagnosis and Disease Monitoring of Acute Promyelocytic Leukemia (APL)</i> ; <b>Methods in Molecular Biology</b> ; Vol. 89; Retinoid Protocols; pages 333-358; (1998).
TS	✓ AJ	Hames, B.D., et al.; <i>Nucleic Acid Hybridisation A Practical Approach</i> ; IRL Press, Oxford; (1985). <i>Table of Contents only</i>
TS	✓ AK	Khrapko, K.R., et al.; <i>An Oligonucleotide Hybridization Approach to DNA Sequencing</i> ; <b>FEBS Letters</b> ; Volume 256, Nos. 1,2; pages 118-122; (1989).
TS	✓ AL	Mangiarotti, Giorgio, et al.; <i>Selection and Analysis of Cloned Developmentally-Regulated Dictyostelium Discoideum Genes by Hybridization-Competition</i> ; <b>Nucleic Acids Research</b> ; Volume 9, No. 4; pages 947-963; (1981).

EXAMINER:

Teresa Stuelcilia

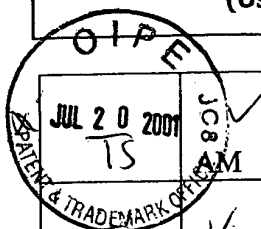
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INFORMATION DISCLOSURE STATEMENTAPPLICANT:  
Chia-Lin WEI, et al.FILING DATE:  
May 23, 2001GROUP:  
To be Assigned 637

(Use several sheets if necessary)



TS	✓	Old, R.W., et al.; <i>Studies in Microbiology; Principles of Gene Manipulation, an Introduction to Genetic Engineering</i> ; Oxford; Blackwell Scientific Publications; Fifth Edition; (1994). <i>Contents only provided.</i>
TS	✓	Riken Genome Exploration Research Group Phase II Team and the Fantom Consortium; <i>Functional Annotation of a Full-Length Mouse cDNA Collection</i> ; <i>Nature</i> ; Vol. 409, Issue No. 6821; pages 685-690; (February 8, 2001).
TS	✓	Rubin, Gerald M., et al.; <i>A Drosophila Complementary DNA Resource</i> ; <i>Science</i> ; Vol. 287, No. 5461; pages 2222-2224; (March 24, 2000).
TS	✓	Sambrook, J., et al.; <i>Molecular Cloning, a Laboratory Manual</i> ; Cold Spring Harbor Laboratory Press; 2 <sup>nd</sup> Ed.; Vols. 1-3; (1989). <i>Table of Contents only.</i>
TS	✓	Sanger, F., et al.; <i>DNA Sequencing with Chain-Terminating Inhibitors</i> ; <i>Proceedings of the National Academy of Sciences of the United States of America</i> ; Vol. 74, No. 12; pages 5463-5467; (December, 1977).
	✓	Sasaki, Yasnory F., et al.; <i>Construction of an Equalized cDNA Library from Human Brain by Semi-Solid Self-Hybridization System</i> ; <i>DNA Research</i> 1; pages 91-96; (1994). ✗
	✓	Schena, Mark, et al.; <i>Parallel Human Genome Analysis: Microarray-Based Expression Monitoring of 1000 Genes</i> ; <i>Proceedings of the National Academy of Sciences of the United States of America</i> ; Vol. 93, No. 20; pages 10614-10619; (October 1, 1996).
	✓	Soares, Marcelo Bento, et al.; <i>Construction and Characterization of a Normalized cDNA Library</i> ; <i>Proceedings of the National Academy of Sciences of the United States of America</i> ; Vol. 91, No. 20; pages 9228-9232; (September 27, 1994).
	✓	Tanaka, Toshihiro et al.; <i>Construction of a Normalized Directionally Cloned cDNA Library from Adult Heart and Analysis of 3040 Clones by Partial Sequencing</i> ; <i>Genomics</i> ; Volume 35; pages 231-235; (1996).
	✓	Wiemann, Stefan, et al.; <i>Toward a Catalog of Human Genes and Proteins: Sequencing and Analysis of 500 Novel Complete Protein Coding Human cDNAs</i> ; <i>Genome Research</i> ; Volume 11, No. 2; pages 422-435 (March 2001).
TS	✓	Wood, William I., et al.; <i>Base Composition-Independent Hybridization in Tetramethylammonium Chloride: A Method for Oligonucleotide Screening of Highly Complex Gene Libraries</i> ; <i>Proceedings of the National Academy of Sciences of the United States of America</i> ; Vol. 82, No. 6; pages 1585-1588; (March 1985).

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T-368 P.005 F-686

<b>LIST OF REFERENCES CITED BY APPLICANT</b> (Use several sheets if necessary) <b>PTO FORM 1449</b>	ATTY. DOCKET NO. 00801.0197.NPUS00	APPLICATION NO. 09/864,637
	APPLICANT Chia-Lin WEI, et al.	
	FILING DATE May 23, 2001	GROUP 1645 637

U.S. PATENT DOCUMENTS							
*EXAMINER INITIAL		DOCUMENT NUMBER	DATE	NAME	CLASS	SUBCLASS	FILING DATE IF APPROPRIATE

FOREIGN PATENT DOCUMENTS								
*EXAMINER INITIAL		DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUBCLASS	TRANSLATION	
							YES	NO

OTHER REFERENCES		
(Including Author, Title, Date, Pertinent Pages, Etc.)		
TS	AX	Nelson et al., "Negative selection: a method for obtaining low-abundance cDNAs using high-density cDNA clone arrays" <i>Genetic Analysis: Biomolecular Engineering</i> 15:209-215 (1999).

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